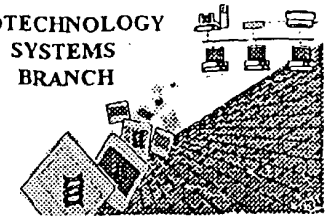
BIOTECHNOLOGY SYSTEMS BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/068,3770
Source: 1600-
Date Processed by STIC: 1/4/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY** or,
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221
Effective 12/13/03: **TELEPHONE: 571-272-2510; FAX: 571-273-0221**

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**>, **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry directly to (EFFECTIVE 12/01/03):**
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. **Federal Express, United Parcel Service, or other delivery service to:** U.S. Patent and Trademark Office, Box Sequence, Room 4B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

ERROR DETECTED | SUGGESTED CORRECTION | SERIAL NUMBER: 09/068,377D

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____ Wrapped Nucleics / Wrapped Aminos — The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2____ Invalid Line Length — The rules require that a line not exceed 72 characters in length. This includes white spaces.

3____ Misaligned Amino Numbering — The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4____ Non-ASCII — The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5____ Variable Length — Sequence(s)____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6____ PatentIn 2.0 "bug" — A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7____ Skipped Sequences (OLD RULES) — Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8____ Skipped Sequences (NEW RULES) — Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9____ Use of n's or Xaa's (NEW RULES) — Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10____ Invalid <213> Response — Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11 J Use of <220> — Sequence(s) 27 missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 07/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12____ PatentIn 2.0 "bug" — Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13____ Misuse of n/Xaa — "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid

1600

RAW SEQUENCE LISTING DATE: 01/04/2004
PATENT APPLICATION: **US/09/068,377D** TIME: 13:06:58

Input Set : **A:\P1066P2.txt**
Output Set: **N:\CRF4\01042004\I068377D.raw**

Does Not Comply
Corrected Diskette Needed

pr 6-7

<110> APPLICANT: Lasky, Laurence A.
Dowbenko, Donald J.
<120> TITLE OF INVENTION: Tyrosine Phosphorylated Cleavage Furrow-Associated Proteins (PSTPIPs)
<130> FILE REFERENCE: P1066P2
<140> CURRENT APPLICATION NUMBER: US 09/068,377D
<141> CURRENT FILING DATE: 1998-05-08
<150> PRIOR APPLICATION NUMBER: PCT/US98/01774
<151> PRIOR FILING DATE: 1998-01-30
<150> PRIOR APPLICATION NUMBER: US 08/938,830
<151> PRIOR FILING DATE: 1997-09-29
<150> PRIOR APPLICATION NUMBER: US 08/798,419
<151> PRIOR FILING DATE: 1997-02-07
<160> NUMBER OF SEQ ID NOS: 76
<210> SEQ ID NO: 1
<211> LENGTH: 415
<212> TYPE: PRT
<213> ORGANISM: Mus Musculus
<400> SEQUENCE: 1

```
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  1               5                  10                  15
Phe Thr Ala His Thr Gly Tyr Glu Val Leu Leu Gln Arg Leu Leu
             20                  25                  30
Asp Gly Arg Lys Met Cys Lys Asp Val Glu Glu Leu Leu Arg Gln
         35                  40                  45
Arg Ala Gln Ala Glu Glu Arg Tyr Gly Lys Glu Leu Val Gln Ile
     50                  55                  60
Ala Arg Lys Ala Gly Gly Gln Thr Glu Met Asn Ser Leu Arg Thr
 65                  70                  75
Ser Phe Asp Ser Leu Lys Gln Gln Thr Glu Asn Val Gly Ser Ala
             80                  85                  90
His Ile Gln Leu Ala Leu Ala Leu Arg Glu Glu Leu Arg Ser Leu
         95                 100                 105
Glu Glu Phe Arg Glu Arg Gln Lys Glu Gln Arg Lys Lys Tyr Glu
    110                 115                 120
Ala Ile Met Asp Arg Val Gln Lys Ser Lys Leu Ser Leu Tyr Lys
125                 130                 135
Lys Thr Met Glu Ser Lys Lys Ala Tyr Asp Gln Lys Cys Arg Asp
140                 145                 150
Ala Asp Asp Ala Glu Gln Ala Phe Glu Arg Val Ser Ala Asn Gly
155                 160                 165
His Gln Lys Gln Val Glu Lys Ser Gln Asn Lys Ala Lys Gln Cys
170                 175                 180
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RAW SEQUENCE LISTING DATE: 01/04/2004
PATENT APPLICATION: US/09/068,377D TIME: 13:06:58

Input Set : A:\P1066P2.txt
Output Set: N:\CRF4\01042004\I068377D.raw

```
Lys Glu Ser Ala Thr Glu Ala Glu Arg Val Tyr Arg Gln Asn Ile
                185                 190                 195
Glu Gln Leu Glu Arg Ala Arg Thr Glu Trp Glu Gln Glu His Arg
                200                 205                 210
Thr Thr Cys Glu Ala Phe Gln Leu Gln Glu Phe Asp Arg Leu Thr
                215                 220                 225
Ile Leu Arg Asn Ala Leu Trp Val His Cys Asn Gln Leu Ser Met
                230                 235                 240
Gln Cys Val Lys Asp Asp Glu Leu Tyr Glu Glu Val Arg Leu Thr
                245                 250                 255
Leu Glu Gly Cys Asp Val Glu Gly Asp Ile Asn Gly Phe Ile Gln
                260                 265                 270
Ser Lys Ser Thr Gly Arg Glu Pro Pro Ala Pro Val Pro Tyr Gln
                275                 280                 285
Asn Tyr Tyr Asp Arg Glu Val Thr Pro Leu Ile Gly Ser Pro Ser
                290                 295                 300
Ile Gln Pro Ser Cys Gly Val Ile Lys Arg Phe Ser Gly Leu Leu
                305                 310                 315
His Gly Ser Pro Lys Thr Thr Pro Ser Ala Pro Ala Ala Ser Thr
                320                 325                 330
 Glu Thr Leu Thr Pro Thr Pro Glu Arg Asn Glu Leu Val Tyr Ala
                 335                 340                 345
 Ser Ile Glu Val Gln Ala Thr Gln Gly Asn Leu Asn Ser Ser Ala
                 350                 355                 360
 Gln Asp Tyr Arg Ala Leu Tyr Asp Tyr Thr Ala Gln Asn Ser Asp
                 365                 370                 375
 Glu Leu Asp Ile Ser Ala Gly Asp Ile Leu Ala Val Ile Leu Glu
                 380                 385                 390
 Gly Glu Asp Gly Trp Trp Thr Val Glu Arg Asn Gly Gln Arg Gly
                 395                 400                 405
 Phe Val Pro Gly Ser Tyr Leu Glu Lys Leu
                 410                 415
<210> SEQ ID NO: 2
<211> LENGTH: 2100
<212> TYPE: DNA
<213> ORGANISM: Mus Musculus
<400> SEQUENCE: 2
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 agaagaagcg gaaggtctcg agcggcgcca attttaatca aagtgggaat 100
 attgctgata gctcattgtc cttcactttc actaacagta gcaacggtcc 150
 gaacctcata acaactcaaa caaattctca agcgctttca caaccaattg 200
 cctcctctaa cgttcatgat aacttcatga ataatgaaat cacggctagt 250
 aaaattgatg atggtaataa ttcaaaacca ctgtcacctg gttggacgga 300
 ccaaactgcg tataacgcgt ttggaatcac tacagggatg tttaatacca 350
 ctacaatgga tgatgtatat aactatctat tcgatgatga agatacccca 400
 ccaaacccaa aaaaagaggg tgggtcgacc cacgcgtccg gctccttcct 450
 catttcgctg ctgattctag ccccaaacaa aacaggttga gccttttttcc 500
 tcctccggca gttgcctctg gcttgtggct gccttctgag cgtttcagac 550
 ggcgccggct gggagtggga gggagggcct gggctagccg cgctgggact 600
```

RAW SEQUENCE LISTING DATE: 01/04/2004
PATENT APPLICATION: US/09/068,377D TIME: 13:06:58

Input Set : A:\P1066P2.txt
Output Set: N:\CRF4\01042004\I068377D.raw

```
148  gggacgtgct cctggctcct ggcccatgct cagccctgct tgaagcagga 650
150  gtgctagcat ttgacacaac gcccttggag gatgatggcc cagctgcagt 700
152  tccgagatgc cttctggtgc agggacttca cggcccacac agggtatgag 750
154  gtgctactgc agaggctgct ggacggcagg aagatgtgca aggatgtgga 800
156  ggagctgctc agacagaggg cccaggcgga ggagaggtac gggaaggagc 850
158  tggtgcagat tgcacgcaag gctggtggcc agacagagat gaattccctg 900
160  aggacctcct ttgactccct gaagcagcaa acagagaatg tgggcagtgc 950
162  acacatccag ctggccctgg ccctgcgtga ggagctgcgg agcctggagg 1000
164  agttccgaga gagacagaaa gagcagcgga agaagtatga ggccatcatg 1050
166  gaccgtgtcc agaagagcaa gttgtcgctc tacaagaaga ccatggagtc 1100
168  caagaaggca tatgaccaga agtgcaggga tgcagatgat gctgagcagg 1150
170  ccttcgagcg tgtgagtgcc aatggccacc agaagcaagt agaaaagagc 1200
172  cagaacaaag ccaagcagtg caaggagtca gccacagagg cagaaagagt 1250
174  gtacaggcaa aatatcgaac aactggagag agcgaggacc gagtgggagc 1300
176  aggagcaccg gactacctgt gaggccttcc agttgcagga gtttgaccgg 1350
178  ctcaccatcc tccgcaatgc cctgtgggtg cactgtaacc agctctccat 1400
180  gcagtgtgtc aaggatgatg agctctatga ggaagtgcgg ctgacccttg 1450
182  agggctgtga tgtggaaggt gacatcaatg gcttcatcca gtccaagagc 1500
184  actggcagag agcccccagc tccggtgcct tatcagaact actatgacag 1550
186  ggaggtgacc ccactgattg gcagccctag catccagccc tcctgcggtg 1600
188  tgataaagag gttctctggg ctgctacatg gaagtcccaa gaccacacct 1650
190  tctgctcctg ctgcttccac agagactctg actcccaccc ctgagcggaa 1700
192  tgagttggtc tacgcatcca tcgaagtgca ggcgacccag ggaaacctta 1750
194  actcatcagc ccaggactac cgggcactct acgactacac tgcacagaat 1800
196  tctgatgagc tggacatttc cgcgggagac atcctggcgg tcatcctgga 1850
198  aggggaggat ggctggtgga ctgtggagcg gaacggacaa cgtggctttg 1900
200  tccctgggtc gtacttggag aagctctgag gaaaggctag cagtctccac 1950
202  atacctccgc cctgactgtg aggtcaggac tgtttctttc catcaccgcc 2000
204  caggcctcac ggggccagaa ccaagcccgg tggtgctggg catgggctgg 2050
206  gtgctggcta ctctcaataa atgtctccca gaaggaaaaa aaaaaaaaaa 2100
208 <210> SEQ ID NO: 3
209 <211> LENGTH: 48
210 <212> TYPE: PRT
211 <213> ORGANISM: Mus Musculus
213 <400> SEQUENCE: 3
214  Leu Tyr Asp Tyr Thr Ala Gln Asn Ser Asp Glu Leu Asp Ile Ser
215    1               5                  10                  15
217  Ala Gly Asp Ile Leu Ala Val Ile Leu Glu Gly Glu Asp Gly Trp
218                   20                  25                  30
220  Trp Thr Val Glu Arg Asn Gly Gln Arg Gly Phe Val Pro Gly Ser
221                   35                  40                  45
223  Tyr Leu Arg
226 <210> SEQ ID NO: 4
227 <211> LENGTH: 50
228 <212> TYPE: PRT
229 <213> ORGANISM: Homo sapien
231 <400> SEQUENCE: 4
232  Leu Tyr Gln Tyr Ile Gly Gln Asp Val Asp Glu Leu Ser Phe Asn
233    1               5                  10                  15
```

RAW SEQUENCE LISTING DATE: 01/04/2004
PATENT APPLICATION: US/09/068,377D TIME: 13:06:58

Input Set : A:\P1066P2.txt
Output Set: N:\CRF4\01042004\I068377D.raw

```
235  Val Asn Glu Val Ile Glu Ile Leu Ile Glu Asp Ser Ser Gly Trp
236                   20                  25                  30
238  Trp Lys Gly Arg Leu His Gly Gln Glu Gly Leu Phe Pro Gly Asn
239                   35                  40                  45
241  Tyr Val Glu Lys Ile
242                   50
244 <210> SEQ ID NO: 5
245 <211> LENGTH: 50
246 <212> TYPE: PRT
247 <213> ORGANISM: Homo sapien
249 <400> SEQUENCE: 5
250  Leu Tyr Asp Tyr Gln Glu Lys Ser Pro Arg Glu Val Thr Met Lys
251    1               5                  10                  15
253  Lys Gly Asp Ile Leu Thr Leu Leu Asn Ser Thr Asn Lys Asp Trp
254                   20                  25                  30
256  Trp Lys Val Glu Val Asn Asp Arg Gln Gly Phe Val Pro Ala Ala
257                   35                  40                  45
259  Tyr Val Lys Lys Leu
260                   50
262 <210> SEQ ID NO: 6
263 <211> LENGTH: 50
264 <212> TYPE: PRT
265 <213> ORGANISM: Homo sapien
267 <400> SEQUENCE: 6
268  Leu Tyr Asp Tyr Gln Gly Glu Gly Ser Asp Glu Leu Ser Phe Asp
269    1               5                  10                  15
271  Pro Asp Asp Ile Ile Thr Asp Ile Glu Met Val Asp Glu Gly Trp
272                   20                  25                  30
274  Trp Arg Gly Gln Cys Arg Gly His Phe Gly Leu Phe Pro Ala Asn
275                   35                  40                  45
277  Tyr Val Lys Leu Leu
278                   50
280 <210> SEQ ID NO: 7
281 <211> LENGTH: 48
282 <212> TYPE: PRT
283 <213> ORGANISM: Homo sapien
285 <400> SEQUENCE: 7
286  Leu Tyr Asp Tyr Gln Ala Ala Gly Asp Asp Glu Ile Ser Phe Asp
287    1               5                  10                  15
289  Pro Asp Asp Ile Ile Thr Asn Ile Glu Met Ile Asp Asp Gly Trp
290                   20                  25                  30
292  Trp Arg Gly Val Cys Lys Gly Arg Tyr Gly Leu Phe Pro Ala Asn
293                   35                  40                  45
295  Tyr Val Glu
298 <210> SEQ ID NO: 8
299 <211> LENGTH: 8
300 <212> TYPE: PRT
301 <213> ORGANISM: Artificial Sequence
303 <220> FEATURE:
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RAW SEQUENCE LISTING DATE: 01/04/2004
PATENT APPLICATION: US/09/068,377D TIME: 13:06:58

Input Set : A:\P1066P2.txt
Output Set: N:\CRF4\01042004\I068377D.raw

304 <223> OTHER INFORMATION: Amino acid epitope tag
306 <400> SEQUENCE: 8
307 Asp Tyr Lys Asp Asp Asp Asp Lys
308 1 5
310 <210> SEQ ID NO: 9
311 <211> LENGTH: 33
312 <212> TYPE: DNA
313 <213> ORGANISM: Artificial Sequence
315 <220> FEATURE:
316 <223> OTHER INFORMATION: Synthetic oligonucleotide probe
318 <400> SEQUENCE: 9
319 cgcggatcca ccatgatggc ccagctgcag ttc 33
321 <210> SEQ ID NO: 10
322 <211> LENGTH: 45
323 <212> TYPE: DNA
324 <213> ORGANISM: Artificial Sequence
326 <220> FEATURE:
327 <223> OTHER INFORMATION: Synthetic oligonucleotide probe
329 <400> SEQUENCE: 10
330 gtacgcgtcg actcacttgt catcgtcgtc cttgtagtcg agctt 45
332 <210> SEQ ID NO: 11
333 <211> LENGTH: 18
334 <212> TYPE: DNA
335 <213> ORGANISM: Artificial Sequence
337 <220> FEATURE:
338 <223> OTHER INFORMATION: Synthetic oligonucleotide probe
340 <400> SEQUENCE: 11
341 tgcctttctc tccacagg 18
343 <210> SEQ ID NO: 12
344 <211> LENGTH: 36
345 <212> TYPE: DNA
346 <213> ORGANISM: Artificial Sequence
348 <220> FEATURE:
349 <223> OTHER INFORMATION: Synthetic oligonucleotide probe
351 <400> SEQUENCE: 12
352 ctccttgagg ttctactagt gggggctggt gtcctg 36
354 <210> SEQ ID NO: 13
355 <211> LENGTH: 39
356 <212> TYPE: DNA
357 <213> ORGANISM: Artificial Sequence
359 <220> FEATURE:
360 <223> OTHER INFORMATION: Synthetic oligonucleotide probe
362 <400> SEQUENCE: 13
363 gcggccgcac tagtatccag tctgtgctcc atctgttac 39
365 <210> SEQ ID NO: 14
366 <211> LENGTH: 17
367 <212> TYPE: DNA
368 <213> ORGANISM: Artificial Sequence
370 <220> FEATURE:

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 01/04/2004
PATENT APPLICATION: **US/09/068,377D** TIME: 13:06:59

Input Set : **A:\P1066P2.txt**
Output Set: **N:\CRF4\01042004\I068377D.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:27; Xaa Pos. 2,3

(see p.7)

<210> SEQ ID NO 27
<211> LENGTH: 4
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Any amino acid
<400> SEQUENCE: 27
Pro Xaa Xaa Pro
1

see Item 11 on Error Summary Sheet

this does not explain "Artificial Sequence"

these need explaining (see p. 6)

VERIFICATION SUMMARY DATE: 01/04/2004
PATENT APPLICATION: **US/09/068,377D** TIME: 13:06:59

Input Set : **A:\P1066P2.txt**
Output Set: **N:\CRF4\01042004\I068377D.raw**

L:711 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:27
L:711 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:27
L:711 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0